

SEQUENCE LISTING

<110> BUTTCHER, Volker et al.

<120> Method for producing alpha-1, 6-branched alpha-1, 4-glucans from sucrose

<130> 0147-0253P

<140>

<141> 2003-11-10

<150> US 09/807,063

<151> 2001-04-09

<160> 34

<170> PatentIn Ver. 2.1

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<211> 2475

<212> DNA

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ccgccacccg caaatggca agcagcttgc catgatggca gacatccgc atg aac cga 178

Met Asn Arg

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Ile Ile Asp Ser Leu Phe Ala Ala Thr His Ser Asp Pro Phe Ala Tyr  
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Leu Gly Arg His Arg Val Asn Asp Glu Arg Glu Ala Val Arg Val Leu  
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Arg Pro Asp Ala His His Ile Asp Ile Ile Asp Arg His Thr Gly Ala  
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Tyr Trp Ile Glu Arg Phe Gly Phe Asp Gly Ile Arg Val Asp Ala Val			
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Gly Asn Arg Ile Ile Val Ile Ser Asn Phe Thr Pro Val Val Arg Glu		
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Gln Arg Asn Ser Ser Leu Lys Asp Ile Asp Ile Ala Arg Glu Asn Asn				
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Thr Arg Ile Leu Asp Ile Tyr Thr Pro Glu Gln Arg Ala Gly Ile Glu	
20 25 30	
Lys Ser Glu Asp Trp Arg Gln Phe Ser Arg Arg Met Asp Thr His Phe	
35 40 45	
Pro Lys Leu Met Asn Glu Leu Asp Ser Val Tyr Gly Asn Asn Glu Ala	
50 55 60	
Leu Leu Pro Met Leu Glu Met Leu Leu Ala Gln Ala Trp Gln Ser Tyr	
65 70 75 80	
Ser Gln Arg Asn Ser Ser Leu Lys Asp Ile Asp Ile Ala Arg Glu Asn	

85

90

95

Asn Pro Asp Trp Ile Leu Ser Asn Lys Gln Val Gly Gly Val Cys Tyr  
100 105 110

Val Asp Leu Phe Ala Gly Asp Leu Lys Gly Leu Lys Asp Lys Ile Pro  
115 120 125

Tyr Phe Gln Glu Leu Gly Leu Thr Tyr Leu His Leu Met Pro Leu Phe  
130 135 140

Lys Cys Pro Glu Gly Lys Ser Asp Gly Gly Tyr Ala Val Ser Ser Tyr  
145 150 155 160

Arg Asp Val Asn Pro Ala Leu Gly Thr Ile Gly Asp Leu Arg Glu Val  
165 170 175

Ile Ala Ala Leu His Glu Ala Gly Ile Ser Ala Val Val Asp Phe Ile  
180 185 190

Phe Asn His Thr Ser Asn Glu His Glu Trp Ala Gln Arg Cys Ala Ala  
195 200 205

Gly Asp Pro Leu Phe Asp Asn Phe Tyr Tyr Ile Phe Pro Asp Arg Arg  
210 215 220

Met Pro Asp Gln Tyr Asp Arg Thr Leu Arg Glu Ile Phe Pro Asp Gln  
225 230 235 240

His Pro Gly Gly Phe Ser Gln Leu Glu Asp Gly Arg Trp Val Trp Thr  
245 250 255

Thr Phe Asn Ser Phe Gln Trp Asp Leu Asn Tyr Ser Asn Pro Trp Val  
260 265 270

Phe Arg Ala Met Ala Gly Glu Met Leu Phe Leu Ala Asn Leu Gly Val  
275 280 285

Asp Ile Leu Arg Met Asp Ala Val Ala Phe Ile Trp Lys Gln Met Gly  
290 295 300

Thr Ser Cys Glu Asn Leu Pro Gln Ala His Ala Leu Ile Arg Ala Phe  
305 310 315 320

Asn Ala Val Met Arg Ile Ala Ala Pro Ala Val Phe Phe Lys Ser Glu  
325 330 335

Ala Ile Val His Pro Asp Gln Val Val Gln Tyr Ile Gly Gln Asp Glu  
340 345 350

Cys Gln Ile Gly Tyr Asn Pro Leu Gln Met Ala Leu Leu Trp Asn Thr  
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Leu Ala Thr Arg Glu Val Asn Leu Leu His Gln Ala Leu Thr Tyr Arg  
370 375 380

His Asn Leu Pro Glu His Thr Ala Trp Val Asn Tyr Val Arg Ser His

385	390	395	400
Asp Asp Ile Gly Trp Thr Phe Ala Asp Glu Asp Ala Ala Tyr Leu Gly			
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Ile Ser Gly Tyr Asp His Arg Gln Phe Leu Asn Arg Phe Phe Val Asn			
420	425	430	
Arg Phe Asp Gly Ser Phe Ala Arg Gly Val Pro Phe Gln Tyr Asn Pro			
435	440	445	
Ser Thr Gly Asp Cys Arg Val Ser Gly Thr Ala Ala Ala Leu Val Gly			
450	455	460	
Leu Ala Gln Asp Asp Pro His Ala Val Asp Arg Ile Lys Leu Leu Tyr			
465	470	475	480
Ser Ile Ala Leu Ser Thr Gly Gly Leu Pro Leu Ile Tyr Leu Gly Asp			
485	490	495	
Glu Val Gly Thr Leu Asn Asp Asp Asp Trp Ser Gln Asp Ser Asn Lys			
500	505	510	
Ser Asp Asp Ser Arg Trp Ala His Arg Pro Arg Tyr Asn Glu Ala Leu			
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Tyr Ala Gln Arg Asn Asp Pro Ser Thr Ala Ala Gly Gln Ile Tyr Gln			
530	535	540	
Gly Leu Arg His Met Ile Ala Val Arg Gln Ser Asn Pro Arg Phe Asp			
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Gly Gly Arg Leu Val Thr Phe Asn Thr Asn Asn Lys His Ile Ile Gly			
565	570	575	
Tyr Ile Arg Asn Asn Ala Leu Leu Ala Phe Gly Asn Phe Ser Glu Tyr			
580	585	590	
Pro Gln Thr Val Thr Ala His Thr Leu Gln Ala Met Pro Phe Lys Ala			
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His Asp Leu Ile Gly Gly Lys Thr Val Ser Leu Asn Gln Asp Leu Thr			
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